In the Claims:

1-123. Canceled.

- 4. 124. (Previously presented) An isolated nucleic acid comprising:
 - (a) a nucleic acid sequence encoding the polypeptide of SEQ ID NO: 377;
 - a nucleic acid sequence encoding the polypeptide of SEQ ID NO: 377,
 lacking its associated signal peptide;
 - (c) the nucleic acid sequence of SEQ ID NO: 376;
 - (d) the full-length coding sequence of the nucleic acid sequence of SEQ ID NO: 376; or
 - (e) the full-length coding sequence of the cDNA deposited under ATCC accession number 203092.
- 5.125. (Previously presented) The isolated nucleic acid of Claim 124 comprising a nucleic acid sequence encoding the polypeptide of SEQ ID NO: 377.
- (Previously presented) The isolated nucleic acid of Claim 124 comprising a nucleic acid sequence encoding the polypeptide of SEQ ID NO: 377, lacking its associated signal peptide.

127-128. Canceled.

- 7, 129. (Previously presented) The isolated nucleic acid of Claim 124 comprising the nucleic acid sequence of SEQ ID NO: 376.
- 8. 130. (Previously presented) The isolated nucleic acid of Claim 124 comprising the full-length coding sequence of the nucleic acid sequence of SEQ ID NO: 376.
- 9, 121. (Previously presented) The isolated nucleic acid of Claim 124 comprising the full-length coding sequence of the cDNA deposited under ATCC accession number 203092.

132-134. Canceled.

10, 125. (Currently amended) A vector comprising the nucleic acid of Claim 124 or 141 139.

- (Previously presented) The vector of Claim 136, wherein said nucleic acid is operably linked to control sequences recognized by a host cell transformed with the vector.
- 12,137. (Previously presented) An isolated host cell comprising the vector of Claim 135.
- 13; (Previously presented) The host cell of Claim 127, wherein said cell is a CHO cell, an E. coli or a yeast cell.

139-140. Canceled.

- 1. (Currently amended) An isolated nucleic acid encoding a polypeptide according to Claim 139 having at least 90% sequence identity to:
 - (a) the amino acid sequence of the polypeptide of SEQ ID NO: 377;
 - (b) the amino acid sequence of the polypeptide of SEQ ID NO: 377, lacking its associated signal peptide;
 - (c) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the nucleic acid sequence of SEQ ID NO: 376; or
 - (d) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 203092; wherein said encoded polypeptide induces chondrocyte redifferentiation.
- (Currently amended) An The isolated nucleic acid encoding a polypeptide according to Claim

 141 139 having at least 95% sequence identity to:
 - (a) the amino acid sequence of the polypeptide of SEQ ID NO: 377;
 - (b) the amino acid sequence of the polypeptide of SEQ ID NO: 377, lacking its associated signal peptide;
 - (c) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the nucleic acid sequence of SEQ ID NO: 376; or
 - (d) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 203092; wherein said encoded polypeptide induces chondrocyte redifferentiation.

- 3.143. (Currently amended) An The isolated nucleic acid encoding a polypeptide according to Claim
 144. 139 having at least 99% sequence identity to:
 - (a) the amino acid sequence of the polypeptide of SEQ ID NO: 377;
 - (b) the amino acid sequence of the polypeptide of SEQ ID NO: 377, lacking its associated signal peptide;
 - (c) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the nucleic acid sequence of SEQ ID NO: 376; or
 - (d) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 203092; wherein said encoded polypeptide induces chondrocyte redifferentiation.